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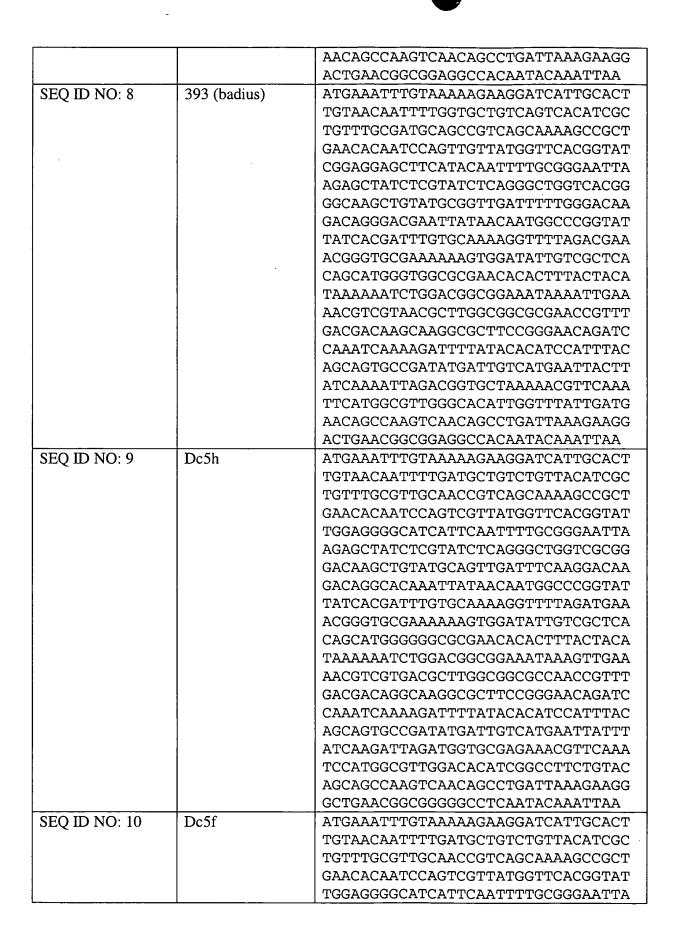
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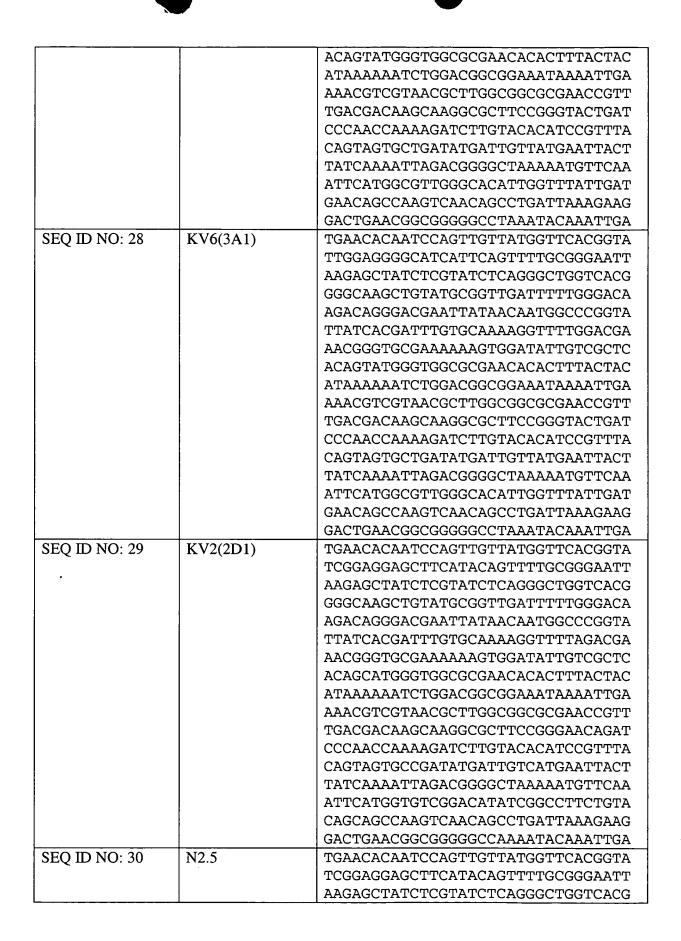
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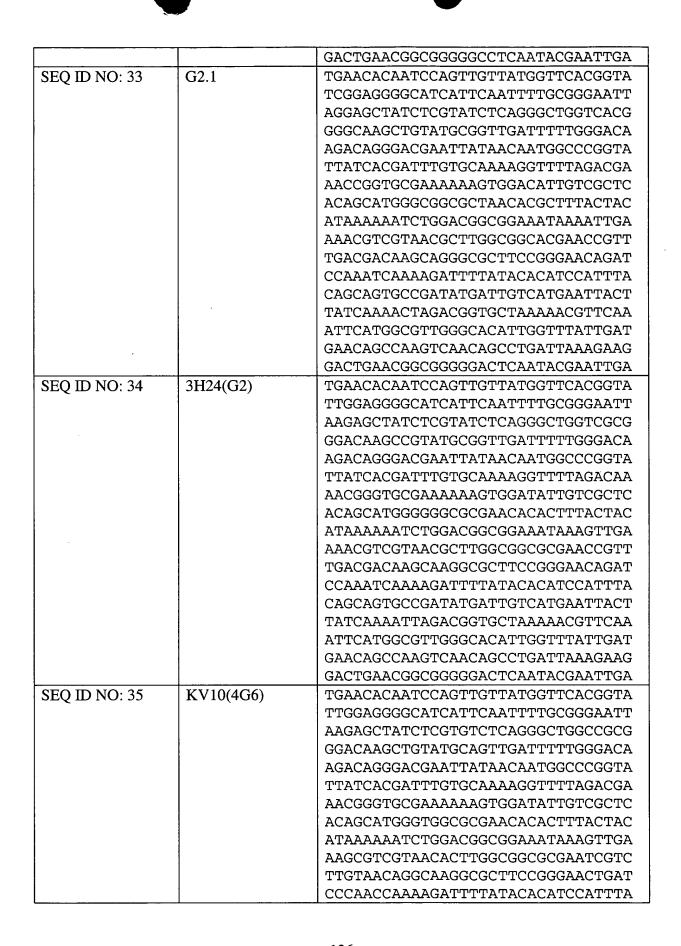
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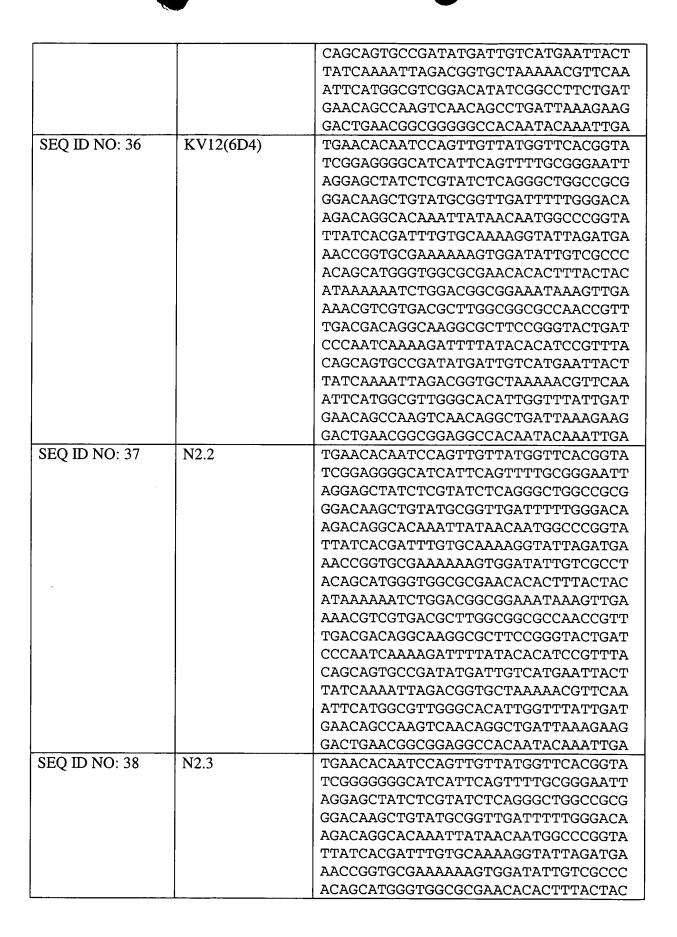
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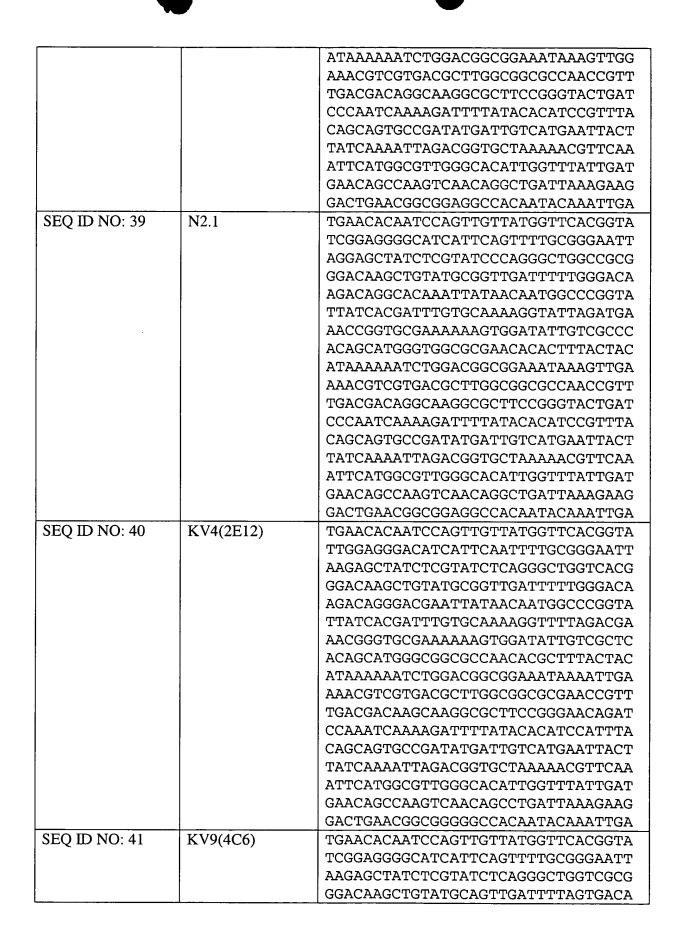
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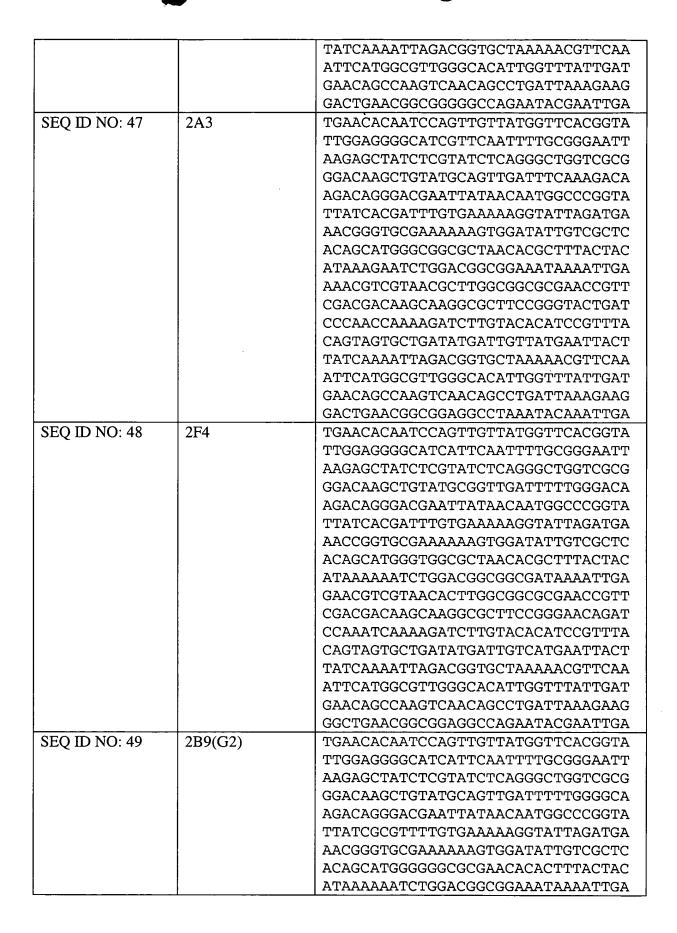






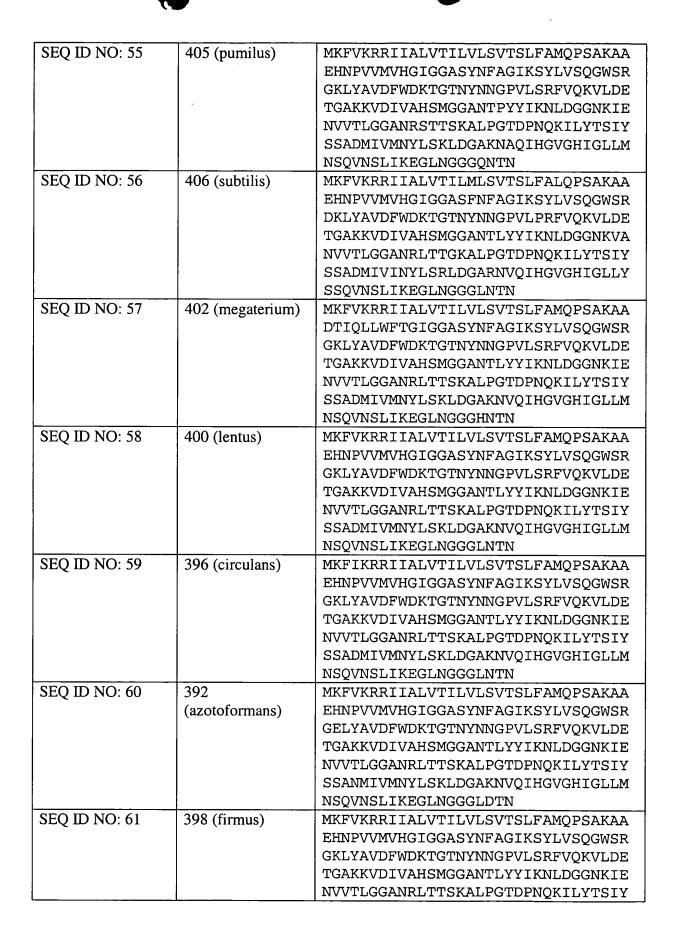
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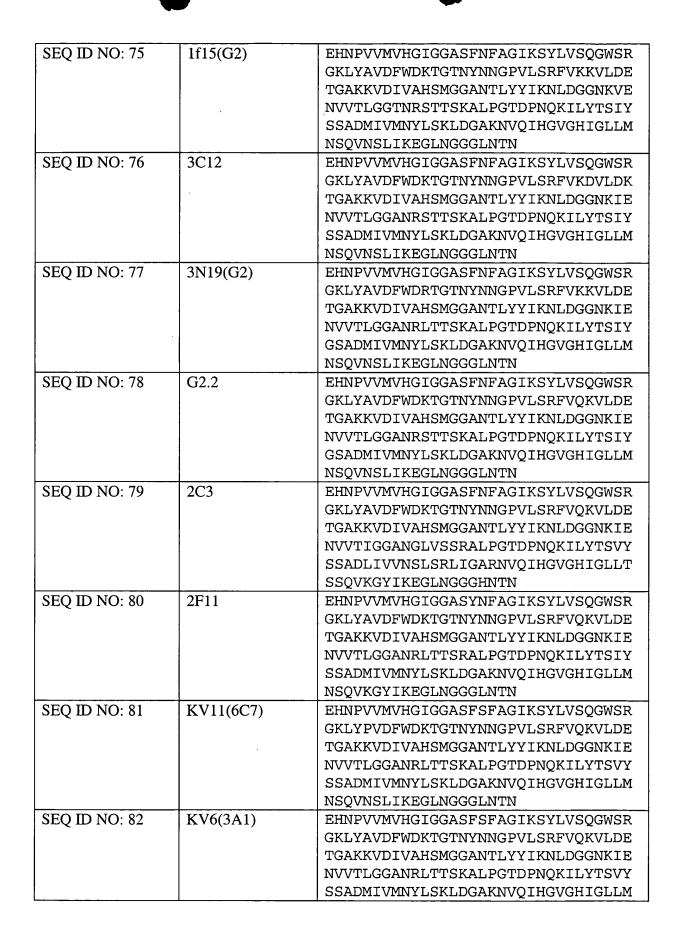


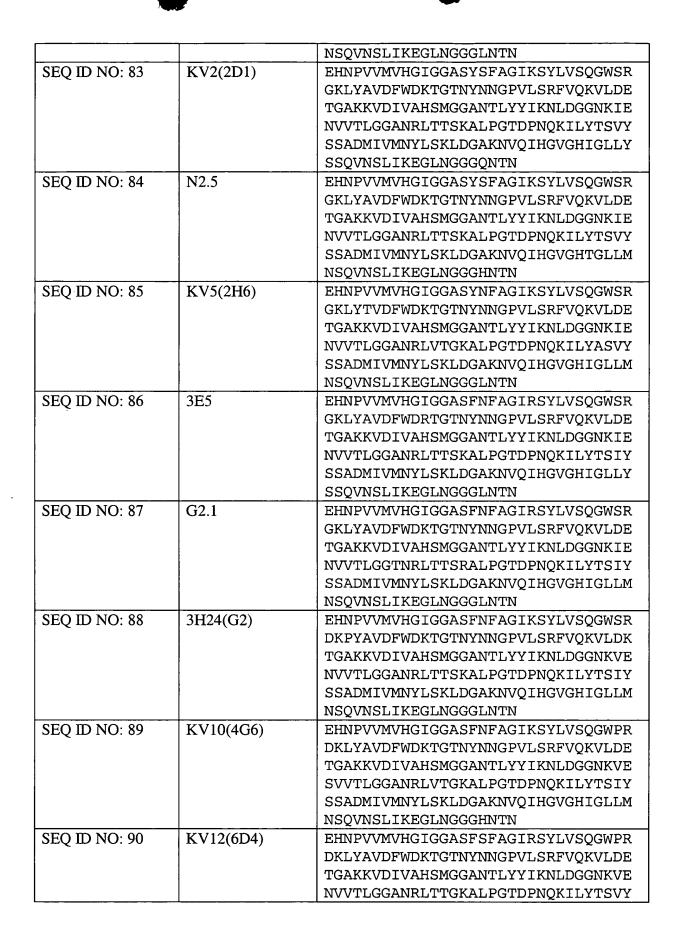


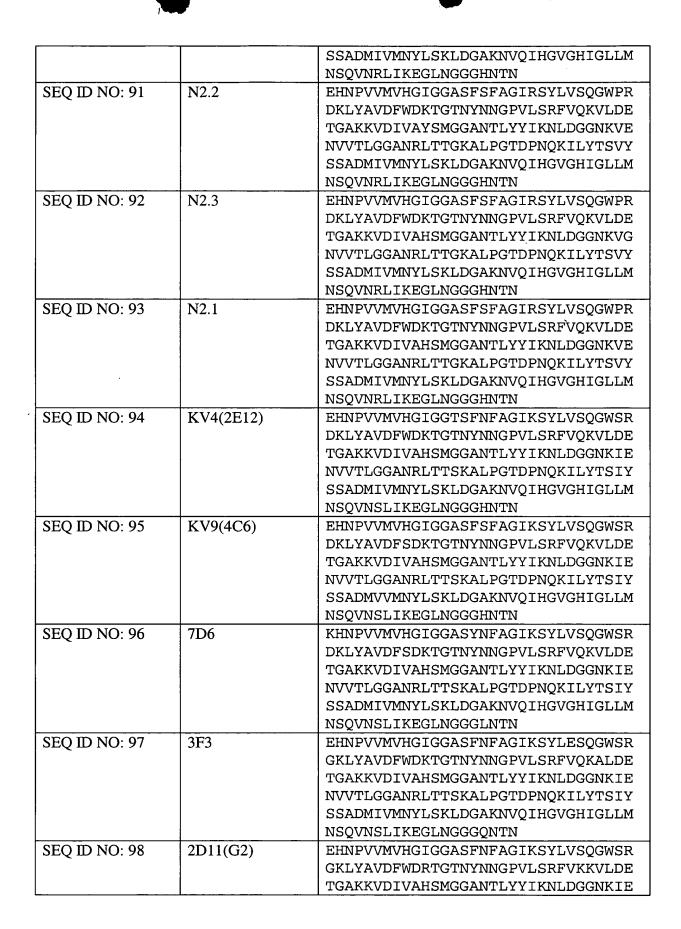


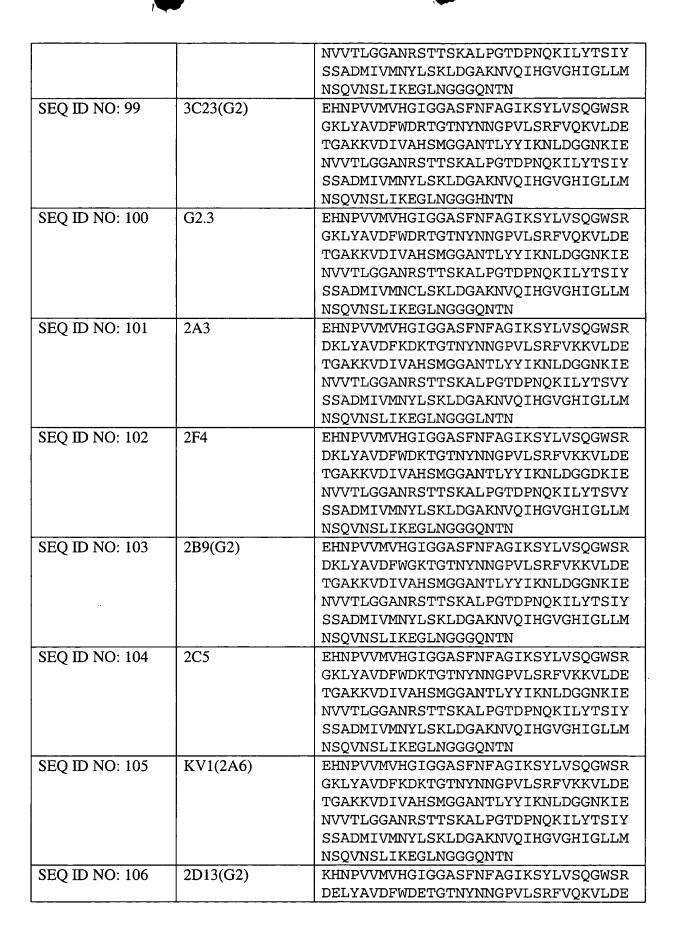
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		SVYSSADLIVVNSLSRLTGARNVLIHGVGHIG
		LLTSSQVKGYIKEGLNGGGLNTN
SEQ ID NO: 74	H2a	MKFVKRRIIALVTILMLSVTSLFALQPSAKAA
		EHNPVVMVHGIGGASFNFAGIKSYLVSQGWSR
		DKLYAVDFRDKTGNNRNNGPRLSKFVKDVLDK
		TGAKKVDIVAHSMGGANTLYYIKNLDGGDKIE
		NVVTIGGANGLVSSRALPGTDPNQKILYTSVY
		KLSRSHCRQQSLSFNWLQETVQIHGVGHIGLL
		TSSQVKGYIKEGLNGGGLNTN
l	<u> </u>	1 - 2 - 2 + 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1













		TGAKKVDIVAHSMGGANTLYYIKNLDGGNKIE
		NVVTLGGANRSTTSKALPGTDPNQKILYTSIY
		SSADMIVMNYLSKLDGAKNVQIHGVGHIGLLM
_		NSQVNSLIKEGLNGGGQNTN
SEQ ID NO: 107	3C8	EHNPVVMVHGIGGASFNFAGIKSYLVSQGWSR
		DKLYAVDFWDKTGTNYNNGPVLSRFVQKVLDE
		TGAKKVDIVAHSMGGANTLYYIKNLDGGNKVE
		NVVTLGGANRSTTSKALPGTDPNQKILYTSIY
		SSADMIVMNYLSKLDGAKNVQIHGVGHIGLLM
		NSQVNSLIKEGLNGGGQNTN
SEQ ID NO: 108	2D5	EHNPVVMVHGIGGASFNFAGIKSYLVSQGWSR
-		GKLYAVDFWDKTGTNYNNGPVLSRFVQKVLDE
		TGAKKVDIVAHSMGGANTLYYIKNLDGGNKIE
		NVVTLGGANRSTTSKALPGTDPNQKILYTSIY
,		SSADMIVMNYLSKLDGAKNVQIHGVGHIGLLM
		NSQVNSLIKEGLNGGGQNTN